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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Thu Oct 25 18:17:00 EDT 2007

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\*\*\*\*\*

Reviewer Comments:

<210> 17

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<212> PRT

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<223> Synthetic peptide

<220>

<221> misc\_feature

<222> (13)..(14)

<223> D-Val

<220>

<221> misc\_feature

<222> (17)..(17)

<223> D-Val

<400> 17

Phe	Ser	Glu	Pro	Glu	Ile	Thr	Leu	Ile	Ile	Phe	Gly	Val	Met	Ala	Gly
1			5					10						15	

Val	Ile	Gly	Thr	Ile	Leu	Leu	Ile	Ser	Tyr	Gly	Ile	Arg	Arg	Leu	Ile
			20					25						30	

One of the above <222> lines indicates Val at locations 13 and 14; Val

is only at location 13. Met is at location 14.

<210> 25  
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<400> 25

Lys Lys Ile Thr Ala Gly Ala Ala Gly Val Ala Ala Gly Val Ala Ala  
1 5 10 15

Ala

One of the above <223> lines indicates "D-Glu" at location 6: "Gly" is  
at location 6.

\*\*\*\*\*

Application No: 10583996

Version No: 2.0

Input Set:

Output Set:

Started: 2007-10-09 10:33:48.696

Finished: 2007-10-09 10:33:50.314

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 618 ms

Total Warnings: 29

Total Errors: 0

No. of SeqIDs Defined: 29

Actual SeqID Count: 29

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W 213	Artificial or Unknown found in <213> in SEQ ID (4)
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Input Set:

Output Set:

Started: 2007-10-09 10:33:48.696  
Finished: 2007-10-09 10:33:50.314  
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 618 ms  
Total Warnings: 29  
Total Errors: 0  
No. of SeqIDs Defined: 29  
Actual SeqID Count: 29

Error code

Error Description

This error has occurred more than 20 times, will not be displayed

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<110> Yeda Research and Development Co. Ltd. at the Weizmann  
Institute of Science

<120> Diastereomeric Peptides Useful As Inhibitors of Membrane Protein  
Assembly

<130> YEDA/038 PCT

<140> 10583996

<141> 2007-10-09

<150> US 60/530,899

<151> 2003-12-22

<160> 29

<170> PatentIn version 3.3

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Tyr Thr Ser Leu Ile His Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln  
1 5 10 15

Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu  
20 25 30

Trp Asn Trp Phe  
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<223> D-Gln

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Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu  
20 25 30

Trp Asn Trp Phe  
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Tyr Thr Ser Leu Ile His Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln  
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Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu  
20 25 30

Trp Asn Trp Phe  
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Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly Ala Ala Gly  
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Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln  
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Leu

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<222> (14)..(14)

<223> D-Ala

<400> 5

Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly Ala Ala Gly  
1 5 10 15

Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln  
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Leu



<210> 6  
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<400> 6

Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly Ala Ala Gly  
1 5 10 15

Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln  
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Leu

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Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly Ala Ala Gly  
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Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln  
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Leu

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Lys Lys Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly Val Ile Gly  
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Thr

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Lys Lys Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly Val Ile Gly  
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Thr Lys Lys

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Thr

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Lys Lys Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly Val Ile Gly  
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Thr Lys Lys

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<222> (15)..(15)

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Thr

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Thr Lys Lys

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Phe Ser Glu Pro Glu Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly  
1 5 10 15

Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg Arg Leu Ile  
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1 5 10 15

Met Ala Gly Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg  
20 25 30

Arg Leu Ile  
35

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Phe Ser Glu Pro Glu Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly  
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Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg Arg Leu Ile  
20 25 30

<210> 20

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Lys Lys Lys Met Val Leu Gly Val Phe Ala Leu Leu Gln Leu Ile Ser  
1 5 10 15

Gly Ser Leu Lys Lys  
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<400> 22

Met Val Leu Gly Val Phe Ala Leu Leu Gln Leu Ile Ser Gly Ser Leu  
1 5 10 15

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Lys Lys Lys Met Val Leu Gly Val Phe Ala Leu Leu Gln Leu Ile Ser  
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Gly Ser Leu Lys Lys  
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Ala Val Gly Ile Gly Ala Leu Phe  
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Lys Lys Ile Thr Ala Gly Ala Ala Gly Val Ala Ala Gly Val Ala Ala  
1 5 10 15

Ala

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Trp Met Glu Trp Asp Arg Glu Ile Asn Asn Tyr Thr Ser Leu Ile His  
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Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Glu  
20 25 30

Leu Leu

<210> 27  
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<400> 27

Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly  
1 5 10

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Phe Phe Gly Ala Val Ile Gly Thr Ile Ala Leu Gly Val Ala Thr Ser  
1 5 10 15

Ala Gln Ile Thr Ala Gly Ile Ala Leu Ala Glu Ala Arg Glu Ala Lys  
20 25 30

Arg

<210> 29  
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<212> PRT  
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<220>  
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<400> 29

Lys Lys Lys Met Val Leu Gly Val Phe Ala Leu Leu Phe Leu Ile Gly  
1 5 10 15

Gly Ser Leu Lys Lys  
20